

GenCore version 4.5
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OM nucleic - nucleic search, using sw model

Run on: May 18, 2002, 02:09:56 ; Search time 8718.72 Seconds
 (without alignments)
 1893.746 Million cell updates/sec

Title: US-09-719-748-1_COPY_98_886

Searched: 789 197056 seqs, 10463268293 residues

Perfect score: 1 tagacatcgagaggact.....ctctcagacaccctggatc 789

Sequence: 1 tagacatcgagaggact.....ctctcagacaccctggatc 789

Scoring table: IDENTITY_NUC

Gapov 10.0, Gapext 1.0

Minimum DB seq length: 0

Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%

Maximum Match 100%

Listing first 45 summaries

Database : GenBank:*

1: gb_ba:*

2: gb_htg:*

3: gb_in:*

4: gb_om:*

5: gb_ov:*

6: gb_pat:*

7: gb_ph:*

8: gb_pi:*

9: gb_pr:*

10: gb_ro:*

11: gb_sts:*

12: gb_sy:*

13: gb_un:*

14: gb_vl:*

15: em_ba:*

16: em_fun:*

17: em_hum:*

18: em_in:*

19: em_mu:*

20: em_om:*

21: em_or:*

22: em_ov:*

23: em_pat:*

24: em_ph:*

25: em_pl:*

26: em_ro:*

27: em_sts:*

28: em_un:*

29: em_vl:*

30: em_htg_hum:*

31: em_htg_inv:*

32: em_htg_other:*

33: em_htg_inv:*

pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

Result No. Score Match Length DB ID Description

8 SUMMARIES

RESULT 1

AF052941 AF052941 LOCUS AF052941 Homo sapiens DAP-kinase related protein 1 mRNA, complete cds. DEFINITION Homo sapiens DAP-kinase related protein 1 mRNA, complete cds. ACCESSION AF052941 VERSION AF052941.1 GI:3560542 KEYWORDS SOURCE human. ORGANISM Homo sapiens Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominoidea; Homo. REFERENCE 1 (bases 1 to 1742) AUTHORS Inbal,B., Shani,G., Cohen,O., Kissil,J.L. and Kimchi,A. TITLE Death-associated protein kinase-related protein 1, a novel Serine/Threonine kinase involved in apoptosis JOURNAL Mol. Cell. Biol. 20 (3), 1044-1054 (2000) MEDLINE 20094983 PUBLMED 1062961 REFERENCE 2 (bases 1 to 1742) AUTHORS Kimchi,A. and Inbal,B. TITLE Direct Submission JOURNAL Submitted (09-MAR-1998) Molecular Genetics, Weizmann Institute of Science, Rehovot 76100, Israel

REFERENCE	1 (bases 1 to 2105)	QY	481	gaatttaagaatattttggacgcggaaatgttgtgtctccagaatgttgcactcgag	540
AUTHORS	Akira, S. and Kawai, T.				
TITLE	Direct Submission				
JOURNAL	Submitted (10-SEP-1997) Shizuo Akira, Hyogo College of Medicine, 663, Japan (E-mail:akirahyo-med.ac.jp, Tel:+81-798-45-6357, Fax:+81-798-46-3164)				
REFERENCE	2 (sites)				
AUTHORS	Kawai, T., Matsumoto, M., Takeda, K., Sanjo, H. and Akira, S.				
TITLE	zip kinase, a novel serine/threonine kinase which mediates apoptosis				
JOURNAL	Mol. Cell. Biol. 18 (3), 1642-1651 (1998)				
MEDLINE	98147805	FEATURES	Location/Qualifiers		
CDS		1. 21.05	/organism="Homo sapiens"		
		2. 14.58	/db_xref="TAXON:9606"		
		3. 14.58	/codon_start=1		
		4. 14.58	/product="zip-kinase"		
		5. 14.58	/protein_id="BAK4955.1"		
		6. 14.58	/db_xref="GI:2911156"		
		7. 14.58	/note="27 a nucleotides"		
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Best Local Similarity	76.6%	Length	2105;		
Matches	604;	Pred.	No. 2.7e-104;		
Conservative	0;	Mismatches	184;	Indels	0;
		Gaps	0;		
polyA_site					
RESULT	12				
AR076189		LOCUS	AR076189	2132 bp	DNA
		DEFINITION	Sequence 3 from patent US 5958748.		
		ACCESION	AR076189		
		VERSION	AR076189.1	61:10002935	
KEYWORDS					
ORGANISM	unknown.	SOURCE			
REFERENCE	1 (bases 1 to 2132)				
AUTHORS	Akira, S. and Kawai, T.				
TITLE	DNA coding for serine/threonine kinase				
JOURNAL	Patent: US 5958748 A 3 28-SEP-1999;				
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BASE COUNT	429 a	ORIGIN	630 c	741 g	332 t
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Best Local Similarity	76.6%	Length	2132;		
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Conservative	0;	Mismatches	184;	Indels	0;
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BASE COUNT	429 a	ORIGIN	630 c	741 g	332 t
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Conservative	0;	Mismatches	184;	Indels	0;
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Best Local Similarity	76.6%	Length	2132;		
Matches	604;	Pred.	No. 2.7e-104;		
Conservative	0;	Mismatches	184;	Indels	0;
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